263LIST83GB.APP SEQUENCE LISTING

```
Commissariat à l'Energie Atomique
<110>
             Centre National de la Recherche Scientifique
             GONDRY Muriel
             GENET Roger
             LAUTRU Sylvie
             PERNODET Jean-Luc
<120> Polynucleotides and polypeptides coded by said polynucleotides
             involved in the synthesis of diketopiperazine derivatives
<130> CGA263/83FR
<140>
<141>
<160> 23
<170> PatentIn Ver. 2.1
<210> 1
<211> 657
<212> DNA
<213> Streptomyces noursei
<400> 1
gtgaggcgcc acccatcgca ttcgccgtac cgcggcgggt gtgaggtgcg cccaaaaaga 60
aggggattga tgttagctca cagttcatct gaatcgccgc cggaatcctt gccggacgcg 120
tǧǧåčggtčc tčaaaåcccg tačcgccgtc čgcaaťtačg cǧåaagagcc ǧgtčǧacǧač 180
gcgctgatcg agcagctgtt ggaggccatg ctcgccgcgc cgaccgcctc caaccggcag 240 gcgtggtcgt tcatggtggt gcgcaggccc gccgcggtcc gcggctgcg cgcgttctcg 300 cccggggtgc tgggaacccc cgccttcttc gtcgtggcct gcgtcgaccg cagtctgacc 360
gacăăcctet cecegaaget ciegeagaag ăteiacgaea ceageaagei etgigiegee 420
atggcggtgg agaacctgct gctcgcggcg cacgcggccg gcctgggcgg atgcccggtg 480 ggcagcttca ggtccgacat cgtcaccagc atgctcggta tcccggaaca catcgagccg 540 atgctcgtgg tcccgatcgg ccgtcccgcg acagccctcg tccctccca gcgccgcgc 600 aagaatgagg tcgtcaacta tgaatcctgg ggaaaccgtg ctgccgccc aactgcg 657
<210> 2
<211> 318
<212> DNA
<213> Streptomyces noursei
<400> 2
atgaatcctg gggaaaccgt gctgccgccc caactgcgtg aggagatcgc gctcctcgcc 60
gtčtatctgč tčágcagcág čcgčggáctc ctggaggágč cágccgacta cggaatttac 120
cgctgtaccg acggggcccg tcgggcgctc caactcctcg acgaacacgg cgggagcacg 180 gcacggctga ccgccgtccg cgagcgtctc gacgaggtca tgttcgcgcc gatgggcgag 240 gaccgggaca tgggcgcgat tctggacgac ctgtgtcgcc aaatggcaga cgctcttccg 300
gaaattgaaa ccccctga
                                                                                                          318
<210> 3
<211> 720
<212> DNA
<213> Streptomyces noursei
<400> 3
atgettgeag gettagttee egegeeggae eaeggaatge gggaagaaat aettggegae 60
cgcagccgat tgatccggca acgcggtgag cacgccctca tcggaatcag tgcgggcaac 120
agttatttca gccagaagaa caccgtcatg ctgctgcaat gggccgggca gcgtttcgag 180 cgcaccgatg tcgtctatgt cgacacccac atcgacgaga tgctgatcgc cgacggccgc 240 agcgcgcagg aggccgagcg gtcggtcaaa cgcacgctca aggatctgcg gcgcagactc 300 cggcgctcgc tggagagcgt gggcgaccac gccgagcggt tccgtgtccg gtccctgtcc 360 gagctccagg agacccctga gtaccgggcc gtacgcgagc gcaccgaccg ggccttcgag 420
```

263LIST83GB.APP gaggacgccg aattcgccac cgcctgcgag gacatggtgc gggccgtggt gatgaaccgg 480 cccggtgacg gcgtcggcat ctccgcggaa cacctgcggg ccggtctgaa ctacgtgctg 540 gccgaggccc cgctcttcgc ggactcgccc ggagtcttct ccgtccctc ctcggtgctc 600 tgctaccaca tcgacacccc gatcacggcg ttcctgtccc ggcgcgagac cggtttccgg 660 gčggccgagg gačaggcgta čgtcgtčgtč aggccčcagg agctggcčga cgčggcctag 720 <210> 4 <211> 834 <212> DNA <213> Streptomyces noursei <400> 4 atgtcatggg gaggacagga cacttgctca tggtgcggaa cggggcccct cggcgaagct 60 gaagacgtag gaagacagca cacgtcgcac gccgggggac ccgtcatgac tcaagccgcc 12 120 accgtcaccg ccaccacgag ccagggcagg gcactcctgc ggagcctgac gccgctgttc 180 gtggacgccg cgatcccgct cggctcgtac ttcctcctcg ccgagggctt cggcatgagc 240 acggtcgccg cgctggcctg gagcagcgtg gtcccggcgc tgcgcacgat ctggggcctg gtccgggagc ggacggtcaa cggcctcgcg ctgctgatcc tcgtcgtcaa cgtggtgggg ctggcgacga gcaccctgac cggcgatgcc cggctgatga tggccaagga cagcggcgtc 300 agcagogtog togggatogo gátoctgoto toggtgogog gcoggogoco gotgátgáco 480 gccggactcc ggccctgggt gaccaaggga agcccggagg ggaacgccgc atgggaccgg 540 ctgtgggcg gcagcgcgg gttccggcaa ctggagcggc gattctcgac ggtctggggg 600 agcgcctgc tgatcgagtg cgtggtcaag gtcgtcgtg cgtacgtcct gccggtgcac 660 accatggtgt ggctgggga ccctggtgcac accatggtgt ggctgggga catggggac catggggac 780 gcgggcggcg gcagcgccga gccgatggag cggatggtca aggccgaggt cggggccgcc 780 ggcgaggccg ccacggcggg gaacgccgag ccggcgccgg ccgccgcggc ctga 834 <210> 5 <211> 3839 <212> DNA <213> Streptomyces noursei <400> 5 ggatccgtcc cgacgggcgg gaaccggtga ggcgccaccc atcgcattcg ccgtaccgcg 60 gcgggtgtga ggtgcgccca aaaagaaggg gattgatgtt agctcacagt tcatctgaat 120 cgccgcgga atccttgccg gacgcgtgga cggtcctcaa aacccgtacc gccgtccgca 180 attacgcgaa agagccggtc gacgacgcgc tgatcgagca gctgttggag gccatgctcg ccgcgccgac cgcctccaac cggcaggcgt ggtcgttcat ggtggtgcgc aggcccgccg 300 cggtccgccg gctgcgcgcg ttctcgcccg gggtgctggg aacccccgcc ttcttcgtcg tggcctgcgt cgaccgcagt ctgaccgaca acctctccc gaagctctcg cagaagatct 420 acgacaccag caagcicigt gicgccatgg cggtggagaa cctgctgctc gcggcgcacg eggeeggeet gggeggatge eeggtgggea getteaggte egaeategte accageatge 540 teggtatece ggaacacate gagecgatge tegtggtee gateggeegt ecegegacag 600 ecetegtee etecagege eggecaaga atgaggtegt caactatgaa teetggggaa 660 accgtgetge egeceaact gegtgaggag ategegetee tegeegteta tetgeteage 720 agcigicgia gactectiga ggageegic gactaeggaa tttaeegetg taeegaegig 780 gčečgteggő egetecaáét cetegaégaa caeggeggga geaeggéaeg getgácegée 840 gtccgcgagc gtctcgacga ggtcatgttc gcgccgatgg gcgaggaccg ggacatgggc 900 gcgattctgg acgacctgtg tcgccaaatg gcagacgctc ttccggaaat tgaaaccccc 960 tgacggctgt ccggggcaac cccaaaagga cttcttagca tgcttgcagg cttagttccc 1020 gcgccggacc acggaatgcg ggaagaaata cttggcgacc gcagccgatt gatccggcaa 1080 cgcggtgagc acgccctcat cggaatcagt gcgggcaaca gttatttcag ccagaagaac 1140 accgtcatgc tgctgcaatg ggccgggcag cgtttcgagc gcaccgatgt cgtctatgtc 1200 gacacccaca tcgacgagat gctgatcgcc gacggccgca gcgcgcagga ggccgagcgg 1260 tcggtcaaac gcacgctcaa ggatctgcgg cgcagactcc ggcgctcgct ggagagcgtg 1320 ggcgaccacg ccgagcggtt ccgtgtccgg tccctgtccg agctccagga gacccctgag 1380 taccgggccg tacgcgagcg caccgaccgg gccttcgagg aggacgccga attcgccacc 1440 gcctgcgagg acatggtgcg ggccgtggtg atgaaccggc ccggtgacgg cgtcggcatc 1500 tccgcggaac acctgcgggc cggtctgaac tacgtgctgg ccgaggcccc gctcttcgcg 1560 gactcgcccg gagtcttctc cgtccctcc tcggtgctct gctaccacat cgacaccccg 1620 atcacggcgt tcctgtcccg gcgcgagacc ggtttccggg cggccgaggg acaggcgtac 1680 gtcgtcgtca ggccccagga gctggccgac gcggcctagt tgggggcgtc cgcgggcgga 1740 cctgcctcc cacccgctc cggtgccggc gccgggcatg acaaatgtca tggggaggac 1800 aggacacttg ctcatggtgc ggaacgggc ccctcggga aggacactt gcacacttg ggaacggag ggaacggaa gaacggaagac 1800 agcacacgtc gcacgccggg ggacccgtca tgactcaagc cgccaccgtc accgccacca 1920 cgagccaggg cagggcactc ctgcggagcc tgacgccgct gttcgtggac gccgcgatcc 1980

```
<210> 6
<211> 219
<212> PRT
<213> Streptomyces noursei
```

Met Arg Arg His Pro Ser His Ser Pro Tyr Arg Gly Gly Cys Glu Val 15 Arg Pro Lys Arg Arg Gly Leu Met Leu Ala His Ser Ser Ser Glu Ser Pro Pro Glu Ser Leu Pro Asp Ala Trp Thr Val Leu Lys Thr Arg Thr Ala Val Arg Asn Tyr Ala Lys Glu Pro Val Asp Asp Ala Leu Ile Glu Gln Leu Leu Glu Ala Met Leu Ala Ala Pro Thr Ala Ser Asn Arg Gln 80 Ala Trp Ser Phe Met Val Val Arg Arg Pro Ala Ala Val Arg Arg Leu Arg Ala Phe Ser Pro Gly Val Leu Gly Thr Pro Ala Phe Phe Val Val Ala Cys Val Asp Asp Arg Ser Leu Thr Asp Asn Leu Ser Pro Lys Leu Ser Gln Lys Ile Tyr Asp Thr Ser Lys Leu Cys Val Ala Met Ala Val Glu Asn Leu Leu Leu Ala Ala His Ala Ala Gly Leu Gly Gly Cys Pro Val 160

Gly Ser Phe Arg Ser Asp Ile Val Thr Ser Met Leu Gly Ile Pro Glu 175

His Ile Glu Pro Met Leu Val Val Pro Ile Gly Arg Pro Ala Thr Ala 180

Leu Val Pro Ser Gln Arg Arg Ala Lys Asn Glu Val Val Asn Tyr Glu 200

Ser Trp Gly Asn Arg Ala Ala Ala Pro Thr Ala 215

<210> 7 <211> 104 <212> PRT <213> Streptomyces noursei

Asn Pro Gly Glu Thr Val Leu Pro Pro Gln Leu Arg Glu Glu Ile Ala Leu Leu Leu Ala Val Tyr Leu Leu Ser Ser Gly Arg Gly Leu Leu Glu Glu Glu Pro Ala Asp Tyr Gly Ile Tyr Arg Cys Thr Asp Gly Ala Arg Arg Ala Leu Gln Leu Leu Asp Glu His Gly Gly Ser Thr Ala Arg Leu Thr Ala Val Arg Glu Arg Leu Asp Glu Val Met Phe Ala Pro Met Gly Glu Asp 65 Arg Asp Met Gly Ala Ile Leu Asp Asp Leu Cys Arg Gln Met Ala Asp Asp Ala Leu Pro Glu Ile Glu Thr Pro

<210> 8 <211> 99 <212> PRT <213> Streptomyces noursei

Leu Ile Gly 35 Ile Ser Ala Gly Asn Ser Tyr Phe Ser Gln Lys Asn Thr Val Met Leu Leu Gln Trp Ala Gly Gln Arg Phe Glu Arg Thr Asp Val Val Tyr Val Asp Thr His Ile Asp Glu Met Leu Ile Ala Asp Gly Arg 80 Ser Ala Gln Glu Ala Glu Arg Ser Val Lys Arg Thr Leu Lys Asp Leu Arg Arg Arg Leu Arg Arg Ser Leu Glu Ser Val Gly Asp His Ala Glu Arg Phe Arg Val Arg Ser Leu Ser Glu Leu Gln Glu Thr Pro Glu Tyr

Arg Ala Val Arg Glu Arg Thr Asp Arg Ala Phe Glu Glu Asp Ala Glu
130

135

140

Phe Ala Thr Ala Cys Glu Asp Met Val Arg Ala Val Val Met Asp Arg

Phe Ala Thr Ala Cys Glu Asp Met Val Arg Ala Val Val Met Asn Arg 145 150 160

Pro Gly Asp Gly Val Gly Ile Ser Ala Glu His Leu Arg Ala Gly Leu 165 170 175

Asn Tyr Val Leu Ala Glu Ala Pro Leu Phe Ala Asp Ser Pro Gly Val 180 185 190

Phe Ser Val Pro Ser Ser Val Leu Cys Tyr His Ile Asp Thr Pro Ile 195 200 205

Thr Ala Phe Leu Ser Arg Arg Glu Thr Gly Phe Arg Ala Ala Glu Gly 210 215 220

Gln Ala Tyr Val Val Val Arg Pro Gln Glu Leu Ala Asp Ala Ala 225 230 235

<210> 10 <211> 277

<212> PRT

<213> Streptomyces noursei

Gly Arg Ala Leu Leu Arg Ser Leu Thr Pro Leu Phe Val Asp Ala Ala 50 60 Ile Pro Leu Gly Ser Tyr Phe Leu Leu Ala Glu Gly Phe Gly Met Ser 65 70 75 80 Thr Val Ala Ala Leu Ala Trp Ser Ser Val Val Pro Ala Leu Arg Thr 85 90 95 Ile Trp Gly Leu Val Arg Glu Arg Thr Val Asn Gly Leu Ala Leu Leu 100 105 110 Ile Leu Val Val Asn Val Val Gly Leu Ala Thr Ser Thr Leu Thr Gly
115 120 125 Asp Ala Arg Leu Met Met Ala Lys Asp Ser Gly Val Ser Ser Val Val 130 135 140 Gly Ile Ala Ile Leu Leu Ser Val Arg Gly Arg Arg Pro Leu Met Thr 145 150 155 160 Ala Gly Leu Arg Pro Trp Val Thr Lys Gly Ser Pro Glu Gly Asn Ala 165 170 175 Ala Trp Asp Arg Leu Trp Ala Arg Ser Ala Arg Phe Arg Gln Leu Glu 180 185 190 Arg Arg Phe Ser Thr Val Trp Gly Ser Ala Leu Leu Ile Glu Cys Val 195 200 205 Val Lys Val Val Gly Ala Tyr Val Leu Pro Val His Thr Met Val Trp 210 215 220 Leu Gly Thr Val Leu Thr Val Val Ala Ile Leu Leu Ala Met Val Val 225 230 235 240 Ala Gly Gly Gly Ser Ala Glu Pro Met Glu Arg Met Val Lys Ala Glu 245 250 255 Val Gly Ala Ala Gly Glu Ala Ala Thr Ala Gly Asn Ala Glu Pro Ala 260 265 270 Pro Ala Ala Ala Ala 275

<210> 11

E - 1

<211> 20 <212> PRT

<213> Streptomyces noursei

<400> 11

Glu Pro Val Asp Asp Ala Leu Ile Glu Gln Leu Leu Glu Ala Met Leu 1 5 10 15

Ala Ala Pro Thr 20

<210> 12

<211> 12 <212> PRT

<213> Streptomyces noursei

<400> 12

Asn Glu Val Val Asn Tyr Glu Xaa Trp Gly Asn Arg 1 5 10

<210> <211> <212> <213>	9	
<400> Gln Al 1	13 a Xaa Ser Phe Met Val Val Arg 5	
<210> <211> <212> <213>	17	
<220> <223>	Description of the artificial sequence:primer	
<400> garccs		17
<210> <211> <212> <213>	17	
<220> <223>	Description of the artificial sequence:primer	
<400> gcgtcg	- -	17
<210> <211> <212> <213>	20	
<220> <223>	Description of the artificial sequence:primer	
<400> aacgar		20
<210> <211> <212> <213>	20	
<220> <223>	Description of the artificial sequence:primer	
<400> tcgtag		20
<210> <211> <212> <213>	20	
<220> <223> I	Description of the artificial sequence:primer	

<400> 18 caggcstggw ssttcatggt	20
<210> 19 <211> 20 <212> DNA <213> Artificial sequence	
<220> <223> Description of the artificial sequence:primer	
<400> 19 accatgaass wccasgcctg	20
<210> 20 <211> 47 <212> DNA <213> Artificial sequence	
<220> <223> Description of the artificial sequence:primer	
<400> 20 cggctgcagg agaagggagc ggacatatgc ttgcaggctt agttccc	47
<210> 21 <211> 42 <212> DNA <213> Artificial sequence	
<220> <223> Description of the artificial sequence:primer	
<400> 21 cggtcccgtg gatccaagct tctaggccgc gtcggccagc tc	42
<210> 22 <211> 36 <212> DNA <213> Artificial sequence	
<220> <223> Description of the artificial sequence:primer	
<400> 22 gagcgggatc ctgcagtgtc atggggagga caggac	36
<210> 23 <211> 41 <212> DNA <213> Artificial sequence	
<220> <223> Description of the artificial sequence:primer	
<400> 23 cgatcacgtg gatccaagct tgccaatcct gtacgcgatt t	41